WHAT IS CLAIMED IS:

- 1. A plant comprising a nucleotide sequence having a modified ARF GAP domain, wherein said plant has decreased organ abscission relative to a plant not having a modified ARF domain.
- 2. The plant of Claim 1, wherein the organ abscission comprises floral abscission.
- 3. The plant of Claim 1, wherein the nucleotide sequence is SEQ ID NO: 3 or SEQ ID NO: 5.
 - 4. The plant of Claim 1, wherein the plant is Arabidopsis thaliana.
- 5. The plant of Claim 1, wherein said decreased organ abscission comprises abolished organ abscission.
- 6. An isolated nucleotide sequence that hybridizes to the complement of the sequence shown in SEQ ID NO: 3 or SEQ ID NO:5 under moderate stringency, wherein expression of said nucleotide sequence in a plant results in reduced or abolished abscission.
- 7. The nucleotide sequence of Claim 6, wherein said nucleotide sequence comprises SEQ ID NO: 3 or SEQ ID NO: 5.
 - 8. A method of preventing organ loss in a plant, comprising: mutating the ARF GAP domain of a gene in a plant; and determining if said mutation results in the prevention of organ loss in said plant.
 - 9. The method of Claim 8, wherein said organ loss is floral organ loss.
- 10. The method of Claim 8, wherein said mutating comprises exposure to ethyl methanesulphonate (EMS).
- 11. The method of Claim 8, wherein said gene comprises the nucleotide sequence of SEQ ID NO: 1.
- 12. The method of Claim 8, wherein said mutating results in said gene expressing a protein that is not full-length.
- 13. The method of Claim 8, wherein said mutating results in said gene expressing an inactive protein.
- 14. The method of Claim 8, wherein said mutating introduces a stop codon into said gene.

15. A isolated polypeptide comprising the amino acid sequence of SEQ ID NO: 4 or SEQ ID NO: 6.